

# illuminaHumanMethylation450kmanifest

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illuminaHumanMethylation450kmanifest

*Manifest object for the illuminaHumanMethylation450k array*

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## Description

This object contains the array design for Illumina's Human Methylation 450k microarray.

## Usage

```
data(illuminaHumanMethylation450kmanifest)
```

## Format

An object of class `illuminaMethylationManifest`

## Details

This package is based on the file `HumanMethylation450_15017482_v.1.2.csv`.

## References

Bibikova, M. et al. (2011) Genomics. High density DNA methylation array with single CpG site resolution. <http://dx.doi.org/10.1016/j.ygeno.2011.07.007>

## See Also

The description of the class in the `minfi` package, [illuminaMethylationManifest-class](#).

## Examples

```
data(illuminaHumanMethylation450kmanifest)
```

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